

<223> Synthetic oligonucleotide probe

<400> 266

cttgactgtc tctgaatctg caccc 25

<210> 267

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic oligonucleotide probe

<400> 267

aagtgggtgga agcctccagt gtgg 24

<210> 268

<211> 52

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic oligonucleotide probe

<400> 268

ccactacggt attagagcaa aagttaaaaa ccatcatggt tcctggagca 50

gc 52

<210> 269

<211> 2747

<212> DNA

<213> Homo sapiens

<400> 269

gcaacctcag cttctagtat ccagactcca gcgccgcccc gggcgcggaac 50

cccaaccccg acccagagct tctccagcgg cggcgcgagcg agcagggctc 100

cccgccttaa cttcctccgc ggggccagc caccttcggg agtccgggtt 150

gcccacctgc aaactctccg cttctgcac ctgccacccc tgagccagcg 200

cgggcccccg agcgagtcac ggccaacgag gggctgcagc tgttgggctt 250

cattctcgcc ttcctgggat ggatcggcgc catcgtcagc actgccctgc 300

cccagtgag gatttactcc tatgccggcg acaacatcgt gaccgccag 350

gccatgtacg aggggctgtg gatgtcctgc gtgtcgcaga gcaccgggca 400

gatccagtgc aaagtctttg actccttgct gaatctgagc agcacattgc 450

aagcaacccg tgccttgatg gtggttgga tcctcctggg agtgatagca 500

atctttgtgg ccaccgttg catgaagtgt atgaagtgt tggaagacga 550

tgaggtgcag aagatgagga tggctgtcat tgggggtgcg atatttcttc 600

ttgcaggctct	ggctattttta	gttgccacag	catggtatg	caatagaatc	650
gttcaagaat	tctatgaccc	tatgaccca	gtcaatgcc	ggtacgaatt	700
tggtcaggct	ctcttcactg	gctgggctgc	tgcttctctc	tgctttctgg	750
gagggtgccct	actttgctgt	tcctgtcccc	gaaaaacaac	ctcttaacca	800
acaccaaggc	cctatccaaa	acctgcacct	tccagcggga	aagactacgt	850
gtgacacaga	ggcaaaagga	gaaaatcatg	ttgaaacaaa	ccgaaaatgg	900
acattgagat	actatcatta	acattaggac	cttagaattt	tgggtattgt	950
aatctgaagt	atggtattac	aaaacaaaca	aacaacaaa	aaacccatgt	1000
gttaaaatac	tcagtgctaa	acatggctta	atcttatttt	atcttctttc	1050
ctcaatatag	gaggggaagt	ttttccattt	gtattactgc	ttcccattga	1100
gtaatcatac	tcaaatgggg	gaaggggtgc	tccttaaata	tatatagata	1150
tgatatata	catgtttttc	tattaaaaat	agacagtaaa	atactattct	1200
cattatgttg	atactagcat	acttaaaata	tctctaaaat	aggtaaattgt	1250
atttaattcc	atattgatga	agatgtttat	tggtatatatt	tctttttcgt	1300
ccttatatac	atatgtaaca	gtcaaatac	atttactctt	cttcattagc	1350
tttgggtgcc	tttgccacaa	gacctagcct	aattttacca	ggatgaattc	1400
tttcaattct	tcattgcgtgc	ccttttcata	tacttatttt	attttttacc	1450
ataatcttat	agcaottgca	tcgttattaa	gcccttattt	gttttgtggt	1500
tcattgggtct	ctatctcctg	aatctaacac	atttcatagc	ctacatttta	1550
gtttctaaag	ccaagaagaa	tttattacaa	atcagaactt	tggaggcaaa	1600
tctttctgca	tgaccaaagt	gataaattcc	tgttgacctt	cccacacaat	1650
ccctgtactc	tgacccatag	cactcttggt	tgctttgaaa	atatttgtcc	1700
aattgagtag	ctgcatgctg	ttcccccagg	tgttgtaaca	caactttatt	1750
gattgaattt	ttaagctact	tattcatagt	tttatatccc	cctaaactac	1800
ctttttgttc	ccatttcctt	aattgtattg	ttttcccaag	tgtaattatc	1850
atgcggtttta	tatcttccta	ataagggtgtg	gtctgtttgt	ctgaacaaag	1900
tgctagactt	tctggagtga	taatctgggtg	acaaatatcc	tctctgtagc	1950
tgtaagcaag	tcacttaatc	tttctacctc	ttttttctat	ctgccaaatt	2000
gagataatga	tacttaacca	gttagaagag	gtagtgtgaa	tattaattag	2050

tttatattac tottattctt tgaacatgaa ctatgcctat gtagtgtctt 2100  
 tatttgctca gctggctgag aactgaaga agtcactgaa caaacctac 2150  
 acacgtacct tcatgtgatt cactgccttc ctctctctac cagtctattt 2200  
 ccaactgaaca aaacctacac acataccttc atgtggttca gtgccttcct 2250  
 ctctctacca gtctatttcc actgaacaaa acctacgcac ataccttcat 2300  
 gtggctcagt gccttcctct ctctaccagt ctatttccat tctttcagct 2350  
 gtgtctgaca tgtttggtgt ctgttccatt ttaacaactg ctcttacttt 2400  
 tccagtctgt acagaatgct atttcacttg agcaagatga tgtaatggaa 2450  
 aggggtgttg cactgggtgtc tggagacctg gatttgagtc ttgggtgctat 2500  
 caatcacogt ctgtgtttga gcaaggcatt tggctgctgt aagcttattg 2550  
 ctctcatctgt aagcgggtgg ttgtaattcc tgatcttccc acctcacagt 2600  
 gatgttggtg ggatccagtg agatagaata catgtaagtg tggttttgta 2650  
 atttaaaaag tgctatacta agggaaagaa ttgaggaatt aactgcatac 2700  
 gttttggtgt tgcttttcaa atgtttgaaa ataaaaaaaa tgттаag 2747

<210> 270

<211> 211

<212> PRT

<213> Homo sapiens

<400> 270

Met	Ala	Asn	Ala	Gly	Leu	Gln	Leu	Leu	Gly	Phe	Ile	Leu	Ala	Phe
1				5					10					15
Leu	Gly	Trp	Ile	Gly	Ala	Ile	Val	Ser	Thr	Ala	Leu	Pro	Gln	Trp
				20					25					30
Arg	Ile	Tyr	Ser	Tyr	Ala	Gly	Asp	Asn	Ile	Val	Thr	Ala	Gln	Ala
				35					40					45
Met	Tyr	Glu	Gly	Leu	Trp	Met	Ser	Cys	Val	Ser	Gln	Ser	Thr	Gly
				50					55					60
Gln	Ile	Gln	Cys	Lys	Val	Phe	Asp	Ser	Leu	Leu	Asn	Leu	Ser	Ser
				65					70					75
Thr	Leu	Gln	Ala	Thr	Arg	Ala	Leu	Met	Val	Val	Gly	Ile	Leu	Leu
				80					85					90
Gly	Val	Ile	Ala	Ile	Phe	Val	Ala	Thr	Val	Gly	Met	Lys	Cys	Met
				95					100					105
Lys	Cys	Leu	Glu	Asp	Asp	Glu	Val	Gln	Lys	Met	Arg	Met	Ala	Val
				110					115					120